WHAT IS CLAIMED IS:

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- 1 1. An isolated sweet taste receptor comprising a T1R3 polypeptide, 2 wherein the T1R3 polypeptide is encoded by a nucleotide sequence that hybridizes under moderately stringent hybridization conditions to a nucleotide sequence encoding an amino 3 4 acid sequence of SEQ ID NO:15, 20, 23, or 25. 1 2. The isolated receptor of claim 1, wherein the T1R3 polypeptide is 2 encoded by a nucleotide sequence that hybridizes under highly stringent hybridization 3 conditions to a nucleotide sequence encoding an amino acid sequence of SEQ ID NO:15, 20, 4 23, or 25. 1 3. The isolated receptor of claim 1, wherein the T1R3 polypeptide has an 2 amino acid sequence of SEQ ID NO:15, 20, 23, or 25. 1 4. The isolated receptor of claim 1, wherein the receptor comprises a 2 T1R3 polypeptide and a heterologous polypeptide. 1 5. The isolated receptor of claim 4, wherein the T1R3 polypeptide and the 2 heterologous polypeptide are non-covalently linked. 1 6. The isolated receptor of claim 4, wherein the T1R3 polypeptide and the 2 heterologous polypeptide are covalently linked. 1 7. The isolated receptor of claim 4, wherein the heterologous polypeptide 2 is a T1R1 polypeptide that is encoded by a nucleotide sequence that hybridizes under 3 moderately stringent hybridization conditions to a nucleotide sequence encoding an amino acid sequence of SEQ ID NO:1, 2, or 3. 4 1 8. The isolated receptor of claim 4, wherein the heterologous polypeptide is a T1R1 polypeptide that is encoded by a nucleotide sequence that hybridizes under highly 2 stringent hybridization conditions to a nucleotide sequence encoding an amino acid sequence 3 4 of SEQ ID NO:1, 2, or 3.
 - 9. The isolated receptor of claim 7, wherein the T1R1 polypeptide has an amino acid sequence of SEQ ID NO:1, 2, or 3.

- 1 10. The isolated receptor of claim 4, wherein the heterologous polypeptide
 2 is a T1R2 polypeptide that is encoded by a nucleotide sequence that hybridizes under
 3 moderately stringent hybridization conditions to a nucleotide sequence encoding an amino
 4 acid sequence of SEQ ID NO:7, 8, or 9.

 1 1. The isolated receptor of claim 4, wherein the heterologous polypeptide
 2 is a T1R2 polypeptide is encoded by a nucleotide sequence that hybridizes under highly
- is a T1R2 polypeptide is encoded by a nucleotide sequence that hybridizes under highly stringent hybridization conditions to a nucleotide sequence encoding an amino acid sequence of SEQ ID NO:7, 8, or 9.
- 1 12. The isolated receptor of claim 10, wherein the T1R2 polypeptide has 2 an amino acid sequence of SEQ ID NO:7, 8, or 9.
- 1 13. The isolated receptor of claim 1, wherein the receptor has G protein 2 coupled receptor activity.
- 1 14. The isolated receptor of claim 1, wherein the receptor specifically 2 binds to antibodies raised against SEQ ID NO: 15, 20, 23, or 25.
- 1 15. An isolated sweet taste receptor comprising a T1R3 polypeptide and a
 2 T1R1 polypeptide, wherein the T1R3 polypeptide is encoded by a nucleotide sequence that
 3 hybridizes under highly stringent hybridization conditions to a nucleotide sequence encoding
 4 an amino acid sequence of SEQ ID NO:15, 20, 23, or 25; and wherein the T1R1 polypeptide
 5 that is encoded by a nucleotide sequence that hybridizes under moderately stringent
 6 hybridization conditions to a nucleotide sequence encoding an amino acid sequence of SEQ
 7 ID NO:1, 2, or 3.
- 1 16. An isolated sweet taste receptor comprising a T1R3 polypeptide and a
 2 T1R2 polypeptide, wherein the T1R3 polypeptide is encoded by a nucleotide sequence that
 3 hybridizes under highly stringent hybridization conditions to a nucleotide sequence encoding
 4 an amino acid sequence of SEQ ID NO:15, 20, 23, or 25; and wherein the T1R2 polypeptide
 5 that is encoded by a nucleotide sequence that hybridizes under moderately stringent
 6 hybridization conditions to a nucleotide sequence encoding an amino acid sequence of SEQ
 7 ID NO:7, 8, or 9.

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17. An antibody that specifically binds to the taste receptor claim 1.

1 18. The antibody of claim 17, wherein the antibody specifically binds to a 2 taste receptor comprising T1R1 and T1R3. 1 19. The antibody of claim 18, wherein the T1R1 polypeptide and the T1R3 2 polypeptide are non-covalently linked. 20. 1 The antibody of claim 18, wherein the T1R1 polypeptide and the T1R3 polypeptide are covalently linked. 2 1 21. The antibody of claim 17, wherein the antibody specifically binds to a 2 taste receptor comprising T1R2 and T1R3. 1 22. The antibody of claim 21, wherein the T1R2 polypeptide and the T1R3 2 polypeptide are non-covalently linked. 1 23. The antibody of claim 21, wherein the T1R2 polypeptide and the T1R3 2 polypeptide are covalently linked. 1 24. A method of identifying a compound that modulates sweet taste signal 2 transduction in taste cells, the method comprising the steps of 3 (i) contacting the compound with a sweet taste receptor comprising a T1R3 4 polypeptide, wherein the T1R3 polypeptide is encoded by a nucleotide sequence that 5 hybridizes under moderately stringent hybridization conditions to a nucleotide sequence 6 encoding an amino acid sequence of SEQ ID NO:15, 20, 23, or 25; and 7 (ii) determining the functional effect of the compound upon the receptor, 8 thereby identifying a compound that modulates sweet signal transduction. 1 The method of claim 24, wherein the T1R3 polypeptide is encoded by 25. 2 a nucleotide sequence that hybridizes under highly stringent hybridization conditions to a 3 nucleotide sequence encoding an amino acid sequence of SEQ ID NO:15, 20, 23, or 25 i 26. The method of claim 24, wherein the receptor comprises a T1R3 2 polypeptide and a heterologous polypeptide.

The method of claim 25, wherein the T1R3 polypeptide and the

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heterologous polypeptide are non-covalently linked.

The method of claim 25, wherein the heterologous polypeptide is a 28. 1 T1R1 polypeptide encoded by a nucleotide sequence that hybridizes under moderately 2 stringent hybridization conditions to a nucleotide sequence encoding an amino acid sequence 3 of SEQ ID NO:1, 2, or 3. 4 The method of claim 25, wherein the heterologous polypeptide is a 29. 1 T1R1 polypeptide encoded by a nucleotide sequence that hybridizes under highly stringent 2 hybridization conditions to a nucleotide sequence encoding an amino acid sequence of SEQ 3 4 ID NO:1, 2, or 3. The method of claim 25, wherein the T1R1 polypeptide has an amino 1 30. 2 acid sequence of SEQ ID NO:1, 2, or 3. The method of claim 25, wherein the heterologous polypeptide is a 31. 1 T1R2 polypeptide encoded by a nucleotide sequence that hybridizes under moderately 2 stringent hybridization conditions to a nucleotide sequence encoding an amino acid sequence 3 of SEQ ID NO:7, 8, or 9. 4 The method of claim 25, wherein the heterologous polypeptide is a 32. 1 T1R2 polypeptide encoded by a nucleotide sequence that hybridizes under highly stringent 2 hybridization conditions to a nucleotide sequence encoding an amino acid sequence of SEQ 3 ID NO:7, 8, or 9. 4 The method of claim 25, wherein the T1R2 polypeptide has an amino 33. 1 acid sequence of SEQ ID NO:6, 7, or 8. 2 The method of claim 24, wherein the receptor is recombinant. 34. 1 The method of claim 24, wherein the receptor has G protein coupled 1 35. receptor activity. 2 The method of claim 24, wherein the functional effect is measured in 36. 1 2 vitro. The method of claim 36, wherein the functional effect is a physical 37. 1

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effect.

| 1 2 | phase. | 38. | The method of claim 36, wherein the receptor is linked to a solid |
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| 1 2 | measuring bind | 39. ding of | The method of claim 36, wherein the functional effect is determined by a compound to the receptor. |
| 1 2 | measuring bind | 40. ding of | The method of claim 39, wherein the functional effect is determined by a compound to the extracellular domain of the receptor. |
| 1 2 | cell membrane | 41 . | The method of claim 24, wherein the receptor is expressed in a cell or |
| 1 2 | effect. | 42. | The method of claim 41, wherein the functional effect is a physical |
| 1 2 | measuring liga | 43. nd bind | The method of claim 42, wherein the functional effect is determined by ling to the receptor. |
| 1 2 | | 44. ding of | The method of claim 43, wherein the functional effect is determined by a compound to the extracellular domain of the receptor. |
| 1 2 | phenotypic eff | 45. ect. | The method of claim 41, wherein the functional effect is a chemical or |
| 1 2 | | 46. nges in | The method of claim 45, wherein the functional effect is determined by intracellular cAMP, IP3, or Ca ²⁺ . |
| 1 | | 47. | The method of claim 41, wherein the cell is a mammalian cell. |
| 1 | | 48. | The method of claim 47, wherein the cell is a human cell. |
| 1 | | 49. | A method of identifying a compound that modulates sweet taste signal |
| 2 | transduction in | taste c | ells, the method comprising the steps of |
| 3 | | (i) con | tacting the compound with cell expressing a sweet taste receptor |
| 4 | comprising a T | 1R3 pc | olypeptide and a T1R2 polypeptide, wherein the T1R3 polypeptide is |
| 5 | encoded by a n | ucleoti | de sequence that hybridizes under highly stringent hybridization |
| 6 | conditions to a | nucleo | tide sequence encoding an amino acid sequence of SEQ ID NO:15, 20, |
| 7 | 23, or 25; and | wherei | n the T1R2 polypeptide that is encoded by a nucleotide sequence that |

| 8 | hybridizes under moderately stringent hybridization conditions to a nucleotide sequence | | |
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| 9 | encoding an amino acid sequence of SEQ ID NO:7, 8, or 9; and | | |
| 10 | (ii) determining the functional effect of the compound upon the receptor, | | |
| 11 | thereby identifying a compound that modulates sweet signal transduction. | | |
| 1 | 50. The method of claim 49, wherein the T1R2 polypeptide and the T1R3 | | |
| 2 | polypeptide are non-covalently linked. | | |
| 1 | 51. The method of claim 49, wherein the T1R2 polypeptide and the T1R3 | | |
| 2 | polypeptide are covalently linked. | | |
| 1 | 52. A method of identifying a compound that modulates sweet taste signal | | |
| 2 | transduction in taste cells, the method comprising the steps of | | |
| 3 | (i) contacting the compound with cell expressing a sweet taste receptor | | |
| 4 | comprising a T1R3 polypeptide and a T1R1 polypeptide, wherein the T1R3 polypeptide is | | |
| 5 | encoded by a nucleotide sequence that hybridizes under highly stringent hybridization | | |
| 6 | conditions to a nucleotide sequence encoding an amino acid sequence of SEQ ID NO:15, 20, | | |
| 7 | 23, or 25; and wherein the T1R1 polypeptide that is encoded by a nucleotide sequence that | | |
| 8 | hybridizes under moderately stringent hybridization conditions to a nucleotide sequence | | |
| 9 | encoding an amino acid sequence of SEQ ID NO:1, 2, or 3; and | | |
| 10 | (ii) determining the functional effect of the compound upon the receptor, | | |
| 11 | thereby identifying a compound that modulates sweet signal transduction. | | |
| 1 | 53. The method of claim 52, wherein the T1R1 polypeptide and the T1R3 | | |
| 2 | polypeptide are non-covalently linked. | | |
| 1 | 54. The method of claim 52, wherein the T1R1 polypeptide and the T1R3 | | |
| 2 | polypeptide are covalently linked. | | |